65 66

-continued

```
<210> SEQ ID NO 96
<211> LENGTH: 21
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: siRNA
      duplex, antisense
<220> FEATURE:
     OTHER INFORMATION: Description of Combined DNA/RNA Molecule:
<223>
      Figure 18
<400> SEQUENCE: 96
ucgaaguaau ccgcguacgt t
                                                                        21
```

The invention claimed is:

- 1. A method of preparing a double-stranded RNA molecule, wherein each RNA strand has a length from 19-25 nucleotides, wherein said RNA molecule is capable of target-specific nucleic acid modifications and wherein at least one strand has a 3'-overhang of 1-5 nucleotides, comprising
 - (a) synthesizing two RNA strands each having a length 25 from 19-25 nucleotides, wherein said RNA strands are capable of forming a double-stranded RNA molecule,
 - (b) combining the synthesized RNA strands under conditions, wherein a double-stranded RNA molecule which mediates target-specific nucleic acid modifica- 30 tions is formed, wherein said double-stranded RNA molecule consists of a single double stranded region and single stranded regions of 1 to 5 nucleotides at the 3' ends of at least one of the strands of said doublestranded RNA molecule.
- 2. The method according to claim 1, wherein the RNA strands are chemically synthesized.
- 3. The method according to claim 1, wherein the RNA strands are enzymatically synthesized.
- 4. The method of claim 1, wherein both strands of said double-stranded RNA each have a 3'-overhang from 1-5 nucleotides.
- 5. The method of claim 1, wherein both strands of said double-stranded RNA each have a 3'-overhang from 1-3 45 double-stranded RNA each have a 3'-overhang from 1-3nucleotides.
- 6. The method of claim 1, wherein both strands of said double-stranded RNA each have a 3'-overhang of 2 nucleotides.
- 7. The method of claim 1, wherein each strand has a 50 length from 20-22 nucleotides.
- 8. The method of claim 7, wherein both strands of said double-stranded RNA each have a 3'-overhang from 1-5 nucleotides.
- 9. The method of claim 7, wherein both strands of said double-stranded RNA each have a 3'-overhang from 1-3 nucleotides.
- 10. The method of claim 7, wherein both strands of said double-stranded RNA each have a 3'-overhang of 2 nucle- 60
- 11. The method of claim 1, wherein the double-stranded RNA comprises at least one sugar-modified ribonucleotide, wherein the 2'-OH group of said sugar-modified ribonucleotide is replaced by a group selected from H, OR, R, halo, 65 SH, SR, NH₂, NHR, N(R)₂ or CN, wherein R is C_1 – C_6 alkyl, alkenyl or alkynyl and halo is F, Cl, Br or I.

- 12. The method of claim 1, wherein the double stranded RNA comprises at least one backbone-modified ribonucleotide containing a phosrhorothioate group.
- 13. A method of preparing a double-stranded RNA molecule, wherein each strand has a length of from 19-25 nucleotides, wherein said RNA molecule is capable of mediating the cleavage of a target mRNA in a mammal and at least one strand has a 3' overhang of 1–3 nucleotides, comprising the steps of:
 - a) selecting a target mammalian mRNA or target gene sequence,
 - (b) synthesizing a first RNA strand having a length from 19-25 nucleotides, wherein said first RNA strand is complementary to contiguous nucleotides in said target mammalian mRNA or said target gene sequence,
 - (c) synthesizing a second RNA strand having a length from 19-25 nucleotides, wherein said second RNA strand is complementary to 16-24 nucleotides from said first RNA strand, and
 - (d) combining the synthesized RNA strands under conditions suitable to form a double stranded RNA molecule, wherein said double stranded RNA molecule consists of a single double stranded region of from 16-24 nucleotides in length and one or two single stranded 3' overhang regions of 1-3 nucleotides in length each.
- 14. The method of claim 13, wherein both strands of said nucleotides.
- 15. The method of claim 13, wherein both strands of said double-stranded RNA each have a 3'-overhang of 2 nucleotides.
- 16. The method of claim 13, wherein each strand has a length from 20-22 nucleotides.
- 17. The method of claim 16, wherein both strands of said double-stranded RNA each have a 3'-overhang from 1-3 nucleotides.
- 18. The method of claim 16, wherein both strands of said double-stranded RNA each have a 3'-overhang of 2 nucleotides.
- 19. The method of claim 13, wherein the double-stranded RNA comprises at least one sugar-modified ribonucleotide, wherein the 2'-OH group of said sugar-modified ribonucleotide is replaced by a group selected from H, OR, R, halo, SH, SR, NH₂, NHR, N(R)₂ or CN, wherein R is C₁-C₆ alkyl, alkenyl or alkynyl and halo is F, Cl, Br or I.
- 20. The method of claim 13, wherein the double stranded RNA comprises at least one backbone-modified ribonucleotide containing a phoshorothioate group.